

Amendment to the specification:

Insert the paper copy of the Sequence Listing filed herewith following the Drawings.

*Please amend the paragraph beginning at page 11, line 16, as follows:*

Figure 1: Secondary VAI RNA Structure of the Adenovirus Serotype 5 (Ad5) (SEQ ID NO:1). A structure of stems and loops formed by the pairing of bases according to Watson and Crick's pairing rules. The central domain is critical to the VA function and the apical stem is also involved in the interaction of VAI RNA with PKR.

*Please amend the paragraph beginning at page 11, line 22, as follows:*

Figure 2: Sequence of the Ad5 VA Region. The DNA sequence shown (SEQ ID NO:2) corresponds to base pairs 10,500 to 11,100 of the adenovirus serotype 5 genome. This sequence contains the VA region (only the strand in the direction of the VA genes is shown). The sequence shown goes from base pairs (bp) -118 in relation to the beginning of the transcription of the VAI gene to 64 bp beyond the termination point of the VAII. The VAI gene (160 bp) from the beginning to the end of the transcription is underlined and italicized. A sequence of 96 bp separates the VAI and VAII encoding sequences. VAII (161 bp) is found after VAI and is shown underlined and bold.